



## RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 10/784,633  
Source: TFUO  
Date Processed by STIC: 7/26/04

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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VERSION 4.2 PROGRAM ACCESSIBLE THROUGH THE U.S. PATENT AND  
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<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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IFWO

## RAW SEQUENCE LISTING

DATE: 07/20/2004

PATENT APPLICATION: US/10/784,633

TIME: 11:10:30

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\07202004\J784633.raw

3 <110> APPLICANT: SHIRAI, Tomoyuki  
 4 ASAMOTO, Makoto  
 5 HOKAIWADO, Naomi  
 7 <120> TITLE OF INVENTION: Carcinogen-hypersensitive rat  
 9 <130> FILE REFERENCE: 671302-3004  
 11 <140> CURRENT APPLICATION NUMBER: 10/784,633  
 12 <141> CURRENT FILING DATE: 2004-02-23  
 14 <150> PRIOR APPLICATION NUMBER: JP P2001-253241  
 15 <151> PRIOR FILING DATE: 2001-08-23  
 17 <150> PRIOR APPLICATION NUMBER: JP 2001-253241  
 18 <151> PRIOR FILING DATE: 2001-08-23  
 20 <160> NUMBER OF SEQ ID NOS: 6  
 22 <170> SOFTWARE: PatentIn Ver. 2.1  
 24 <210> SEQ ID NO: 1  
 25 <211> LENGTH: 1485  
 26 <212> TYPE: DNA  
 27 <213> ORGANISM: Rattus norvegicus  
 29 <220> FEATURE:  
 30 <221> NAME/KEY: CDS  
 31 <222> LOCATION: (32)..(883)  
 33 <400> SEQUENCE: 1  
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 35 Met Asn Trp Thr Gly Leu Tyr  
 36 1 5  
 38 acc ttg ctc agt ggc gtg aat cgg cat tct aca gcc att ggc cga gta 100  
 39 Thr Leu Leu Ser Gly Val Asn Arg His Ser Thr Ala Ile Gly Arg Val  
 40 10 15 20  
 42 tgg ctg tcc gtc atc ttt atc ttc aga atc atg gtg ctg gtg gtg gct 148  
 43 Trp Leu Ser Val Ile Phe Ile Phe Arg Ile Met Val Leu Val Val Ala  
 44 25 30 35  
 46 gca gag agc gtg tgg ggt gat gag aag tct tct ttc atc tgt aac acc 196  
 47 Ala Glu Ser Val Trp Gly Asp Glu Lys Ser Ser Phe Ile Cys Asn Thr  
 48 40 45 50 55  
 50 ctc cag ccg ggc tgt aac agc gtc tgc tat gac cat ttt ttc ccc atc 244  
 51 Leu Gln Pro Gly Cys Asn Ser Val Cys Tyr Asp His Phe Phe Pro Ile  
 52 60 65 70  
 54 tcc cat gtg cgc ctg tgg tcc ctg caa ctc atc ttg gtt tcc acc cca 292  
 55 Ser His Val Arg Leu Trp Ser Leu Gln Leu Ile Leu Val Ser Thr Pro  
 56 75 80 85  
 58 gct ctc ctc gtg gca atg cac gtg gct cac caa caa cac ata gaa aag 340  
 59 Ala Leu Leu Val Ala Met His Val Ala His Gln Gln His Ile Glu Lys  
 60 90 95 100  
 62 aaa atg cta cgg ctt gag ggg cac ggg gac ccc ctt cac ctg gaa gag 388

Does Not Comply  
Corrected Diskette Needed  
(Pg. 3-4)

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64      105      110      115
66 gta aag agg cac aag gtg cac atc tca ggg aca ctg tgg tgg acc tat 436
67 Val Lys Arg His Lys Val His Ile Ser Gly Thr Leu Trp Trp Thr Tyr
68 120      125      130      135
70 gtc atc agt gtg gtg ttc cgg ctg ctg ttt gag gct gtc ttc atg tat 484
71 Val Ile Ser Val Val Phe Arg Leu Leu Phe Glu Ala Val Phe Met Tyr
72      140      145      150
74 gtc ttc tat ctg ctc tac ccg ggc tat gcc atg gtg cgg ctg gtc aag 532
75 Val Phe Tyr Leu Leu Tyr Pro Gly Tyr Ala Met Val Arg Leu Val Lys
76      155      160      165
78 tgt gag gcc ttc ccc tgc ccc aac acg gtg gac tgc ttc gtg tcc cgc 580
79 Cys Glu Ala Phe Pro Cys Pro Asn Thr Val Asp Cys Phe Val Ser Arg
80      170      175      180
82 ccc act gag aaa acc gtc ttc act gtc ttt atg ctc gcc gcc tcc ggc 628
83 Pro Thr Glu Lys Thr Val Phe Thr Val Phe Met Leu Ala Ala Ser Gly
84      185      190      195
86 atc tgc att atc ctc aac gtg gcg gag gtg gtg tac ctc atc atc cgg 676
87 Ile Cys Ile Ile Leu Asn Val Ala Glu Val Tyr Leu Ile Ile Arg
88 200      205      210      215
90 gcc tgt gcc cgc cgt gct cag cgc cgc tcc aat ccg ccc tcc cgc aag 724
91 Ala Cys Ala Arg Arg Ala Gln Arg Arg Ser Asn Pro Pro Ser Arg Lys
92      220      225      230
94 ggc tgc ggc ttc ggc cac cgc ctc tca cct gaa tac aag cag aat gag 772
95 Gly Ser Gly Phe Gly His Arg Leu Ser Pro Glu Tyr Lys Gln Asn Glu
96      235      240      245
98 atc aac aag ctg ctg agc gag cag gat ggc tct ctg aaa gac ata ctg 820
99 Ile Asn Lys Leu Leu Ser Glu Gln Asp Gly Ser Leu Lys Asp Ile Leu
100      250      255      260
102 cgc cgc agt cct ggc act ggg gcc ggg ctg gct gag aag agc gac cga 868
103 Arg Arg Ser Pro Gly Thr Gly Ala Gly Leu Ala Glu Lys Ser Asp Arg
104      265      270      275
106 tgc tca gcc tgc tga tgccgagtac caggcaacct cccatccaac ccctccctca 923
107 Cys Ser Ala Cys
108 280
110 cccacccag gcctgccct ccttctccta tgcgtggtgag caggcctctg cctcctaggg 983
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124 aggacacatc taggggtggg gagttctgga gggagaagca ggcagataaa tcagagtggg 1403
126 ggttggtcag ggctgcccc agtccccagt tcccaaggcc tctctctctg aaaatgttac 1463
128 acattaaca ggattttaca gt 1485
131 <210> SEQ ID NO: 2
132 <211> LENGTH: 283
133 <212> TYPE: PRT
134 <213> ORGANISM: Rattus norvegicus

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136 <400> SEQUENCE: 2
137 Met Asn Trp Thr Gly Leu Tyr Thr Leu Leu Ser Gly Val Asn Arg His
138 1 5 10 15
139 Ser Thr Ala Ile Gly Arg Val Trp Leu Ser Val Ile Phe Ile Phe Arg
140 20 25 30
141 Ile Met Val Leu Val Val Ala Ala Glu Ser Val Trp Gly Asp Glu Lys
142 35 40 45
143 Ser Ser Phe Ile Cys Asn Thr Leu Gln Pro Gly Cys Asn Ser Val Cys
144 50 55 60
145 Tyr Asp His Phe Phe Pro Ile Ser His Val Arg Leu Trp Ser Leu Gln
146 65 70 75 80
147 Leu Ile Leu Val Ser Thr Pro Ala Leu Leu Val Ala Met His Val Ala
148 85 90 95
149 His Gln Gln His Ile Glu Lys Lys Met Leu Arg Leu Glu Gly His Gly
150 100 105 110
151 Asp Pro Leu His Leu Glu Glu Val Lys Arg His Lys Val His Ile Ser
152 115 120 125
153 Gly Thr Leu Trp Trp Thr Tyr Val Ile Ser Val Val Phe Arg Leu Leu
154 130 135 140
155 Phe Glu Ala Val Phe Met Tyr Val Phe Tyr Leu Leu Tyr Pro Gly Tyr
156 145 150 155 160
157 Ala Met Val Arg Leu Val Lys Cys Glu Ala Phe Pro Cys Pro Asn Thr
158 165 170 175
159 Val Asp Cys Phe Val Ser Arg Pro Thr Glu Lys Thr Val Phe Thr Val
160 180 185 190
161 Phe Met Leu Ala Ala Ser Gly Ile Cys Ile Ile Leu Asn Val Ala Glu
162 195 200 205
163 Val Val Tyr Leu Ile Ile Arg Ala Cys Ala Arg Arg Ala Gln Arg Arg
164 210 215 220
165 Ser Asn Pro Pro Ser Arg Lys Gly Ser Gly Phe Gly His Arg Leu Ser
166 225 230 235 240
167 Pro Glu Tyr Lys Gln Asn Glu Ile Asn Lys Leu Leu Ser Glu Gln Asp
168 245 250 255
169 Gly Ser Leu Lys Asp Ile Leu Arg Arg Ser Pro Gly Thr Gly Ala Gly
170 260 265 270
171 Leu Ala Glu Lys Ser Asp Arg Cys Ser Ala Cys
172 275 280

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176 &lt;210&gt; SEQ ID NO: 3

177 &lt;211&gt; LENGTH: 21

178 &lt;212&gt; TYPE: DNA

179 &lt;213&gt; ORGANISM: Artificial Sequence

181 &lt;220&gt; FEATURE:

182 &lt;223&gt; OTHER INFORMATION: Insertion Sequence

184 &lt;400&gt; SEQUENCE: 3

185 catcatcacc atcaccattg a

188 &lt;210&gt; SEQ ID NO: 4

189 &lt;211&gt; LENGTH: 20

190 &lt;212&gt; TYPE: DNA

191 &lt;213&gt; ORGANISM: Artificial Sequence

Invalid Response

What is the source of  
Genetic material?

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TIME: 11:10:30

Input Set : A:\PTO.LM.txt

Output Set : N:\CRF4\07202004\J784633.raw

193 &lt;220&gt; FEATURE:

194 &lt;223&gt; OTHER INFORMATION: P1

196 &lt;400&gt; SEQUENCE: 4

197 aacgtggcgc aggtggtgta

200 &lt;210&gt; SEQ ID NO: 5

201 &lt;211&gt; LENGTH: 20

202 &lt;212&gt; TYPE: DNA

203 &lt;213&gt; ORGANISM: Artificial Sequence

205 &lt;220&gt; FEATURE:

206 &lt;223&gt; OTHER INFORMATION: P2

208 &lt;400&gt; SEQUENCE: 5

209 atggtgatgg tgatgatggc

212 &lt;210&gt; SEQ ID NO: 6

213 &lt;211&gt; LENGTH: 21

214 &lt;212&gt; TYPE: DNA

215 &lt;213&gt; ORGANISM: Artificial Sequence

217 &lt;220&gt; FEATURE:

218 &lt;223&gt; OTHER INFORMATION: P3

220 &lt;400&gt; SEQUENCE: 6

221 gggaagggtt gatggagtaa t

20

20

21

What is the source of  
genetic material?

same error

same error

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/784,633

DATE: 07/20/2004

TIME: 11:10:31

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\07202004\J784633.raw